

ENTERED

OIPE

1636

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/975,132A

DATE: 07/10/2002

TIME: 10:51:53

Input Set : A:\GC636-2-SEQLIST.TXT

Output Set: N:\CRF3\07102002\I975132A.raw

4 <110> APPLICANT: Kolkman, Marc
6 <120> TITLE OF INVENTION: Enhanced Secretion of a Polypeptide by a
7 Microorganism
9 <130> FILE REFERENCE: GC636-2
11 <140> CURRENT APPLICATION NUMBER: US 09/975,132A
12 <141> CURRENT FILING DATE: 2001-10-09
14 <150> PRIOR APPLICATION NUMBER: US 60/239,531
15 <151> PRIOR FILING DATE: 2000-10-10
17 <160> NUMBER OF SEQ ID NOS: 30
19 <170> SOFTWARE: FastSEQ for Windows Version 4.0
21 <210> SEQ ID NO: 1
22 <211> LENGTH: 15
23 <212> TYPE: PRT
24 <213> ORGANISM: Artificial Sequence
26 <220> FEATURE:
27 <223> OTHER INFORMATION: synthetic peptide tag
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31 1 5 10 15
33 <210> SEQ ID NO: 2
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35 <212> TYPE: PRT
36 <213> ORGANISM: Artificial Sequence
38 <220> FEATURE:
39 <223> OTHER INFORMATION: synthetic peptide tag
41 <400> SEQUENCE: 2
42 Ala Gly Lys Thr Asn Ser Phe Asn Gln Asn Val Ala Leu Asp Asp
43 1 5 10 15
45 <210> SEQ ID NO: 3
46 <211> LENGTH: 11
47 <212> TYPE: PRT
48 <213> ORGANISM: Artificial Sequence
50 <220> FEATURE:
51 <223> OTHER INFORMATION: synthetic peptide tag
53 <400> SEQUENCE: 3
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55 1 5 10
57 <210> SEQ ID NO: 4
58 <211> LENGTH: 15
59 <212> TYPE: PRT
60 <213> ORGANISM: Artificial Sequence
62 <220> FEATURE:
63 <223> OTHER INFORMATION: synthetic peptide tag

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65 <400> SEQUENCE: 4
66 Ala Gly Lys Thr Asn Ser Phe Asn Gln Asn Val Ala Leu Lys Lys
67 1 5 10 15
69 <210> SEQ ID NO: 5
70 <211> LENGTH: 14
71 <212> TYPE: PRT
72 <213> ORGANISM: Artificial Sequence
74 <220> FEATURE:
75 <223> OTHER INFORMATION: synthetic peptide tag
77 <400> SEQUENCE: 5
78 Gly Lys Thr Asn Ser Phe Asn Gln Asn Val Ala Leu Ala Ala
79 1 5 10
81 <210> SEQ ID NO: 6
82 <211> LENGTH: 14
83 <212> TYPE: PRT
84 <213> ORGANISM: Artificial Sequence
86 <220> FEATURE:
87 <223> OTHER INFORMATION: synthetic peptide tag
89 <400> SEQUENCE: 6
90 Gly Lys Thr Asn Ser Phe Asn Gln Asn Val Ala Leu Asp Asp
91 1 5 10
93 <210> SEQ ID NO: 7
94 <211> LENGTH: 32
95 <212> TYPE: DNA
96 <213> ORGANISM: Artificial Sequence
98 <220> FEATURE:
99 <223> OTHER INFORMATION: primer
101 <400> SEQUENCE: 7
102 gtcgacctcg agacccaag cttggcgtaa tc 32
104 <210> SEQ ID NO: 8
105 <211> LENGTH: 60
106 <212> TYPE: DNA
107 <213> ORGANISM: Artificial Sequence
109 <220> FEATURE:
110 <223> OTHER INFORMATION: primer
112 <400> SEQUENCE: 8
113 gtcgacctcg agcggcagaa tctttttttg attctgccgc aaagtcgtct gttgagcctg 60
115 <210> SEQ ID NO: 9
116 <211> LENGTH: 71
117 <212> TYPE: DNA
118 <213> ORGANISM: Artificial Sequence
120 <220> FEATURE:
121 <223> OTHER INFORMATION: primer
123 <400> SEQUENCE: 9
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125 ttggcgtaat c 71
127 <210> SEQ ID NO: 10
128 <211> LENGTH: 81
129 <212> TYPE: DNA

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130 <213> ORGANISM: Artificial Sequence
132 <220> FEATURE:
133 <223> OTHER INFORMATION: primer
135 <400> SEQUENCE: 10
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137 gctcaaagtc gtctgttgag c      81
139 <210> SEQ ID NO: 11
140 <211> LENGTH: 81
141 <212> TYPE: DNA
142 <213> ORGANISM: Artificial Sequence
144 <220> FEATURE:
145 <223> OTHER INFORMATION: primer
147 <400> SEQUENCE: 11
148 cttctactcg agtcagtcgt ctaatgctac gttttggtta aaactgttag ttttgccctgc      60
149 gctcaaagtc gtctgttgag c      81
151 <210> SEQ ID NO: 12
152 <211> LENGTH: 69
153 <212> TYPE: DNA
154 <213> ORGANISM: Artificial Sequence
156 <220> FEATURE:
157 <223> OTHER INFORMATION: primer
159 <400> SEQUENCE: 12
160 cttctactcg agtcaagctg ctaaagcgta gttttcgctcg tttgctgcgc tcaaagtcgt      60
161 ctgttgagc      69
163 <210> SEQ ID NO: 13
164 <211> LENGTH: 21
165 <212> TYPE: DNA
166 <213> ORGANISM: Artificial Sequence
168 <220> FEATURE:
169 <223> OTHER INFORMATION: primer
171 <400> SEQUENCE: 13
172 cagctccgtc tgaggaaaaa g      21
174 <210> SEQ ID NO: 14
175 <211> LENGTH: 21
176 <212> TYPE: DNA
177 <213> ORGANISM: Artificial Sequence
179 <220> FEATURE:
180 <223> OTHER INFORMATION: primer
182 <400> SEQUENCE: 14
183 cgaagtgggc gatttcttcc g      21
185 <210> SEQ ID NO: 15
186 <211> LENGTH: 40
187 <212> TYPE: DNA
188 <213> ORGANISM: Artificial Sequence
190 <220> FEATURE:
191 <223> OTHER INFORMATION: primer
193 <400> SEQUENCE: 15
194 ttctaaaagc ttagtgcttg attcgaaaat caggcctgtg      40
196 <210> SEQ ID NO: 16

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199 <213> ORGANISM: Artificial Sequence
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205 gagctcgtcg cgcttattag tcgtotaatg ctacgttttg gttaa      45
207 <210> SEQ ID NO: 17
208 <211> LENGTH: 45
209 <212> TYPE: DNA
210 <213> ORGANISM: Artificial Sequence
212 <220> FEATURE:
213 <223> OTHER INFORMATION: primer
215 <400> SEQUENCE: 17
216 ttaaccaaaa cgtagcatta gacgactaat aagcgcagcg agctc      45
218 <210> SEQ ID NO: 18
219 <211> LENGTH: 40
220 <212> TYPE: DNA
221 <213> ORGANISM: Artificial Sequence
223 <220> FEATURE:
224 <223> OTHER INFORMATION: primer
226 <400> SEQUENCE: 18
227 cctccgtgca tgcttcctct tatttattga cagaaatctg      40
229 <210> SEQ ID NO: 19
230 <211> LENGTH: 40
231 <212> TYPE: DNA
232 <213> ORGANISM: Artificial Sequence
234 <220> FEATURE:
235 <223> OTHER INFORMATION: primer
237 <400> SEQUENCE: 19
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240 <210> SEQ ID NO: 20
241 <211> LENGTH: 40
242 <212> TYPE: DNA
243 <213> ORGANISM: Artificial Sequence
245 <220> FEATURE:
246 <223> OTHER INFORMATION: primer
248 <400> SEQUENCE: 20
249 tgtatatgta aatttcagat catcataaat atctgctatt      40
251 <210> SEQ ID NO: 21
252 <211> LENGTH: 31
253 <212> TYPE: DNA
254 <213> ORGANISM: Artificial Sequence
256 <220> FEATURE:
257 <223> OTHER INFORMATION: primer
259 <400> SEQUENCE: 21
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262 <210> SEQ ID NO: 22
263 <211> LENGTH: 30

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264 <212> TYPE: DNA
265 <213> ORGANISM: Artificial Sequence
267 <220> FEATURE:
268 <223> OTHER INFORMATION: primer
270 <400> SEQUENCE: 22
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273 <210> SEQ ID NO: 23
274 <211> LENGTH: 24
275 <212> TYPE: DNA
276 <213> ORGANISM: Artificial Sequence
278 <220> FEATURE:
279 <223> OTHER INFORMATION: primer
281 <400> SEQUENCE: 23
282 acgttacgga ttgacaggg atgg 24
284 <210> SEQ ID NO: 24
285 <211> LENGTH: 22
286 <212> TYPE: DNA
287 <213> ORGANISM: Artificial Sequence
289 <220> FEATURE:
290 <223> OTHER INFORMATION: primer
292 <400> SEQUENCE: 24
293 gagtcgaacc cacgtccaga aa 22
295 <210> SEQ ID NO: 25
296 <211> LENGTH: 152
297 <212> TYPE: PRT
298 <213> ORGANISM: Bacillus subtilis
300 <400> SEQUENCE: 25
301 Met Ser Arg Leu Pro Val Leu Leu Leu Leu Gln Leu Leu Val Arg Pro
302 1 5 10 15
303 Gly Leu Gln Ala Pro Met Thr Gln Thr Thr Pro Leu Lys Thr Ser Trp
304 20 25 30
305 Val Asn Cys Ser Asn Met Ile Asp Glu Ile Ile Thr His Leu Lys Gln
306 35 40 45
307 Pro Pro Leu Pro Leu Leu Asp Phe Asn Asn Leu Asn Gly Glu Asp Gln
308 50 55 60
309 Asp Ile Leu Met Glu Asn Asn Leu Arg Arg Pro Asn Leu Glu Ala Phe
310 65 70 75 80
311 Asn Arg Ala Val Lys Ser Leu Gln Asn Ala Ser Ala Ile Glu Ser Ile
312 85 90 95
313 Leu Lys Asn Leu Leu Pro Cys Leu Pro Leu Ala Thr Ala Ala Pro Thr
314 100 105 110
315 Arg His Pro Ile His Ile Lys Asp Gly Asp Trp Asn Glu Phe Arg Arg
316 115 120 125
317 Lys Leu Thr Phe Tyr Leu Lys Thr Leu Glu Asn Ala Gln Ala Gln Gln
318 130 135 140
319 Thr Thr Leu Ser Leu Ala Ile Phe
320 145 150
322 <210> SEQ ID NO: 26
323 <211> LENGTH: 158

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VERIFICATION SUMMARY

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